



SEQUENCE LISTING

Pepinsky, et al.

<120> HYDROPHOBICALLY-MODIFIED PROTEIN COMPOSITIONS AND METHODS

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<140> 09/579,680

<141> 2000-05-26

<150> PCT/US98/25676

<151> 1998-12-13

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<170> PatentIn Ver. 2.1

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<212> PRT

<213> human

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1 5 10 15Leu Val Pro Leu Ala Tyr Lys Gln Phe Ser Pro Asn Val Pro Glu Lys
20 25 30Thr Leu Gly Ala Ser Gly Arg Tyr Glu Gly Lys Ile Ala Arg Ser Ser
35 40 45Glu Arg Phe Lys Glu Leu Thr Pro Asn Tyr Asn Pro Asp Ile Ile Phe
50 55 60Lys Asp Glu Glu Asn Thr Gly Ala Asp Arg Leu Met Thr Gln Arg Cys
65 70 75 80Lys Asp Arg Leu Asn Ser Leu Ala Ile Ser Val Met Asn Gln Trp Pro
85 90 95Gly Val Lys Leu Arg Val Thr Glu Gly Trp Asp Glu Asp Gly His His
100 105 110Ser Glu Glu Ser Leu His Tyr Glu Gly Arg Ala Val Asp Ile Thr Thr
115 120 125Ser Asp Arg Asp Arg Asn Lys Tyr Gly Leu Leu Ala Arg Leu Ala Val
130 135 140Glu Ala Gly Phe Asp Trp Val Tyr Tyr Glu Ser Lys Ala His Val His
145 150 155 160Cys Ser Val Lys Ser Glu His Ser Ala Ala Ala Lys Thr Gly Gly
165 170 175

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 Thr Pro Leu Ala Tyr Lys Gln Phe Ile Pro Asn Val Ala Glu Lys Thr
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 Leu Gly Ala Ser Gly Arg Tyr Glu Gly Lys Ile Ser Arg Asn Ser Glu
 35 40 45
 Arg Phe Lys Glu Leu Thr Pro Asn Tyr Asn Pro Asp Ile Ile Phe Lys
 50 55 60
 Asp Glu Glu Asn Thr Gly Ala Asp Arg Leu Met Thr Gln Arg Cys Lys
 65 70 75 80
 Asp Lys Leu Asn Ala Leu Ala Ile Ser Val Met Asn Gln Trp Pro Gly
 85 90 95
 Val Lys Leu Arg Val Thr Glu Gly Trp Asp Glu Asp Gly His His Ser
 100 105 110
 Glu Glu Ser Leu His Tyr Glu Gly Arg Ala Val Asp Ile Thr Thr Ser
 115 120 125
 Asp Arg Asp Arg Ser Lys Tyr Gly Met Leu Ala Arg Leu Ala Val Glu
 130 135 140
 Ala Gly Phe Asp Trp Val Tyr Tyr Glu Ser Lys Ala His Ile His Cys
 145 150 155 160
 Ser Val Lys Ala Glu Asn Ser Val Ala Ala Lys Ser Gly Gly
 165 170

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 Gln Leu Val Pro Leu Leu Tyr Lys Gln Phe Val Pro Gly Val Pro Glu
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 Arg Thr Leu Gly Ala Ser Gly Pro Ala Glu Gly Arg Val Ala Arg Gly
 35 40 45

Ser	Glu	Arg	Phe	Arg	Asp	Leu	Val	Pro	Asn	Tyr	Asn	Pro	Asp	Ile	Ile
50					55						60				
Phe	Lys	Asp	Glu	Glu	Asn	Ser	Gly	Ala	Asp	Arg	Leu	Met	Thr	Glu	Arg
65				70						75				80	
Cys	Lys	Glu	Arg	Val	Asn	Ala	Leu	Ala	Ile	Ala	Val	Met	Asn	Met	Trp
				85					90				95		
Pro	Gly	Val	Arg	Leu	Arg	Val	Thr	Glu	Gly	Trp	Asp	Glu	Asp	Gly	His
				100				105				110			
His	Ala	Gln	Asp	Ser	Leu	His	Tyr	Glu	Gly	Arg	Ala	Leu	Asp	Ile	Thr
				115			120				125				
Thr	Ser	Asp	Arg	Asp	Arg	Asn	Lys	Tyr	Gly	Leu	Leu	Ala	Arg	Leu	Ala
				130			135				140				
Val	Glu	Ala	Gly	Phe	Asp	Trp	Val	Tyr	Tyr	Glu	Ser	Arg	Asn	His	Val
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His	Val	Ser	Val	Lys	Ala	Asp	Asn	Ser	Leu	Ala	Val	Arg	Ala	Gly	Gly
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Cys Gly Pro Gly Arg Xaa Xaa Xaa Xaa Xaa Arg Arg Xaa Xaa Xaa Lys
 1 5 10 15

Xaa Leu Xaa Pro Leu Xaa Tyr Lys Gln Phe Xaa Pro Xaa Val Xaa Glu
 20 25 30

Lys Thr Leu Gly Ala Ser Gly Arg Xaa Glu Gly Lys Xaa Xaa Arg Xaa
 35 40 45

Ser Glu Arg Phe Lys Xaa Leu Xaa Pro Asn Tyr Asn Pro Asp Ile Ile
 50 55 60

Phe Lys Asp Glu Glu Asn Xaa Gly Ala Asp Arg Leu Met Thr Xaa Arg
 65 70 75 80

Cys Lys Xaa Xaa Xaa Asn Ser Leu Ala Ile Xaa Val Met Asn Xaa Trp
 85 90 95

Pro Gly Val Lys Leu Arg Val Thr Glu Gly Trp Asp Glu Asp Gly His
 100 105 110

His Xaa Xaa Xaa Ser Leu His Tyr Glu Gly Arg Ala Val Asp Ile Thr
 115 120 125

Thr Ser Asp Arg Asp Arg Xaa Lys Tyr Gly Xaa Leu Ala Arg Leu Ala
 130 135 140

Val Glu Ala Gly Phe Asp Trp Val Tyr Tyr Glu Ser Xaa Xaa His Xaa
 145 150 155 160

His Xaa Ser Val Lys Xaa Xaa Xaa Ser Xaa Ala Ala Xaa Xaa Gly Gly
 165 170 175